## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

/0/519,838
PCT/10
1/19/05

## ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 01/19/2005

PATENT APPLICATION: US/10/519,838

TIME: 11:49:06

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\01192005\J519838.raw

2 <110> APPLICANT: Mitrani, Eduardo N.

4 <120> TITLE OF INVENTION: METHOD AND DEVICE FOR INDUCING BIOLOGICAL PROCESSES BY

## MICRO-ORGANS

6 <130> FILE REFERENCE: 26463

C--> 8 <140> CURRENT APPLICATION NUMBER: US/10/519,838

C--> 8 <141> CURRENT FILING DATE: 2005-01-12

8 <160> NUMBER OF SEQ ID NOS: 8

10 <170> SOFTWARE: PatentIn Ver. 3.1

## ERRORED SEQUENCES

Does Not Compty
Corrected Diskette Neede:

81 <210> SEQ ID NO: 8

82 <211> LENGTH: 18

83 <212> TYPE: DNA

84 <213> ORGANISM: Artificial sequence

W--> 85 <220> FEATURE:

86 <223> OTHER INFORMATION: Single strand DNA oligonucleotide

88 <400> SEQUENCE: 8

C--> 89 ccgccttggc ttgtcaca 18

E--> 96(1

VERIFICATION SUMMARY DATE: 01/19/2005
PATENT APPLICATION: US/10/519,838 TIME: 11:49:07

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\01192005\J519838.raw

L:8 M:270 C: Current Application Number differs, Replaced Current Application No L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:16 M:283 W: Missing Blank Line separator, <220> field identifier L:18 M:283 W: Missing Blank Line separator, <400> field identifier L:19 M:112 C: (48) String data converted to lower case, L:25 M:283 W: Missing Blank Line separator, <220> field identifier L:29 M:112 C: (48) String data converted to lower case, L:35 M:283 W: Missing Blank Line separator, <220> field identifier L:39 M:112 C: (48) String data converted to lower case, L:45 M:283 W: Missing Blank Line separator, <220> field identifier L:49 M:112 C: (48) String data converted to lower case, L:55 M:283 W: Missing Blank Line separator, <220> field identifier L:59 M:112 C: (48) String data converted to lower case, L:65 M:283 W: Missing Blank Line separator, <220> field identifier L:69 M:112 C: (48) String data converted to lower case, L:75 M:283 W: Missing Blank Line separator, <220> field identifier L:79 M:112 C: (48) String data converted to lower case, L:85 M:283 W: Missing Blank Line separator, <220> field identifier L:89 M:112 C: (48) String data converted to lower case, L:96 M:254 E: No. of Bases conflict, this line has no nucleotides.